



Ala Ile Ala Leu Ser Ala Leu Ser Leu Ser Phe Gly Leu Gln Ala Ser  
 -15 -10 -5 1

5 gag tta cca caa att cca cca cag cag gtg aac aac acc atg tac cag 144  
 Glu Leu Pro Gln Ile Pro Pro Gln Gln Val Asn Asn Thr Met Tyr Gln  
 5 10 15

10 gca ttt tat tgg gat gcc tac cct ggc ctt tgg gcc aat tta ccg gcc 192  
 Ala Phe Tyr Trp Asp Ala Tyr Pro Gly Leu Trp Ala Asn Leu Pro Ala  
 10 20 25 30

15 atg gcg gcc cct ttg gcc gag cgt ggc att acc tcg atg tgg ttg ccg 240  
 Met Ala Ala Pro Leu Ala Glu Arg Gly Ile Thr Ser Met Trp Leu Pro  
 35 40 45

20 ccc gcc gcc aaa ggc atg aat ggt act ttc agt gtc ggt tac gat gta 288  
 Pro Ala Ala Lys Gly Met Asn Gly Thr Phe Ser Val Gly Tyr Asp Val  
 50 55 60 65

25 tac gat ttc tgg gat ctg ggc gag ttt aac caa aaa ggc acc acc gcc 336  
 Tyr Asp Phe Trp Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Thr Ala  
 70 75 80

30 acc cgt tac ggt act cgt cag cag ctg caa caa gca ctg agt gct ctg 384  
 Thr Arg Tyr Gly Thr Arg Gln Gln Leu Gln Gln Ala Leu Ser Ala Leu  
 85 90 95

35 gac caa ctg ggt att cag gcc tat ttt gat gtg gtg ttt aac cac cgc 432  
 Asp Gln Leu Gly Ile Gln Ala Tyr Phe Asp Val Val Phe Asn His Arg  
 100 105 110

40 atg ggc gcc gat gca cag gag aat att cct ggc ttt ggc ctg gcc tgg 480  
 Met Gly Ala Asp Ala Gln Glu Asn Ile Pro Gly Phe Gly Leu Ala Trp  
 115 120 125

45 acc gag tat cat ctg caa ggt cgt cag gcg cat tat acc cag caa aac 528  
 Thr Glu Tyr His Leu Gln Gly Arg Gln Ala His Tyr Thr Gln Gln Asn  
 130 135 140 145

	tgg ggc tac ttg tgg cac gac ttt gac tgg aac tgg acc gcg ttt aat	576
	Trp Gly Tyr Leu Trp His Asp Phe Asp Trp Asn Trp Thr Ala Phe Asn	
	150 155 160	
5	ggc tcc gac aat cag ctc tac ccc ggc aaa tgg tgg ggc aat acc acc ttc	624
	Gly Ser Asp Asn Gln Leu Tyr Pro Gly Lys Trp Trp Gly Asn Thr Phe	
	165 170 175	
10	cac ttc cct tat ttg atg ggt gag gat gtc gat tac aac cgc ttt gaa	672
	His Phe Pro Tyr Leu Met Gly Glu Asp Val Asp Tyr Asn Arg Phe Glu	
	180 185 190	
15	gtg cag cag gaa atg aaa gcc tgg ggc gag tgg atc atc aac agc gtt	720
	Val Gln Gln Glu Met Lys Ala Trp Gly Glu Trp Ile Ile Asn Ser Val	
	195 200 205	
20	ggc ttt agc ggc ttt cgg atg gat gcc atc gcc cat gtc gat acc gat	768
	Gly Phe Ser Gly Phe Arg Met Asp Ala Ile Ala His Val Asp Thr Asp	
	210 215 220 225	
	ttt acc cgt gac tgg atc aat cac gtg cag tgg gcc acc agt gag gat	816
	Phe Thr Arg Asp Trp Ile Asn His Val Gln Trp Ala Thr Ser Glu Asp	
	230 235 240	
25	gtg ttc ttt gtc gct gaa gcc tgg gtc agt gat atc aac ggc tat ctg	864
	Val Phe Phe Val Ala Glu Ala Trp Val Ser Asp Ile Asn Gly Tyr Leu	
	245 250 255	
30	gat gca gtc aat acg ccg cat ttg cgc gct ttt gat ttc aat ttg cgc	912
	Asp Ala Val Asn Thr Pro His Leu Arg Ala Phe Asp Phe Asn Leu Arg	
	260 265 270	
35	gaa gac ttc gtt gct tta agc agc ggt agc aaa gac atg cgt tgg tgg	960
	Glu Asp Phe Val Ala Leu Ser Ser Gly Ser Lys Asp Met Arg Trp Trp	
	275 280 285	
	ggc ggt ctg gtc aat agc cag cac cgt gat cgg gcg gtc act ttt gtc	1008
	Gly Gly Leu Val Asn Ser Gln His Arg Asp Arg Ala Val Thr Phe Val	
	290 295 300 305	

	gat aac cac gat acc agc cgg gcc ggc aac cct tat ggc atg ccg cag		1056
	Asp Asn His Asp Thr Ser Arg Ala Gly Asn Pro Tyr Gly Met Pro Gln		
	310	315	320
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	gtg atc aac tac aag aac cag gcc tac gct tac att ctg ttg cgt gag		1104
	Val Ile Asn Tyr Lys Asn Gln Ala Tyr Ala Tyr Ile Leu Leu Arg Glu		
	325	330	335
10			
	cat ggg gtg ccg act gtg ttt gcc cgc gat tac gac gaa ttt ggc atg		1152
	His Gly Val Pro Thr Val Phe Ala Arg Asp Tyr Asp Glu Phe Gly Met		
	340	345	350
15			
	gcg cca acg ctg gat aaa ttg att gag gcg cgc cgc tac ttt gct tat		1200
	Ala Pro Thr Leu Asp Lys Leu Ile Glu Ala Arg Arg Tyr Phe Ala Tyr		
	355	360	365
20			
	ggt cct ggc cat gag tac tcc ggc aat acc gag gcc gtc tac gcc tat		1248
	Gly Pro Gly His Glu Tyr Ser Gly Asn Thr Glu Ala Val Tyr Ala Tyr		
	370	375	380
	385		
	gtg cgc gaa ggg ctt agc act gtg ccg ggt acc ggt ctg gtg atg ctg		1296
	Val Arg Glu Gly Leu Ser Thr Val Pro Gly Thr Gly Leu Val Met Leu		
	390	395	400
25			
	ata tcg ggt cga aac tgg ggt cag cag tcg ttc acc atc aac agc		1344
	Ile Ser Gly Arg Asn Trp Gly Gly Gln Gln Ser Phe Thr Ile Asn Ser		
	405	410	415
30			
	cac cag ccg aat acc acc ttt tac gat tat acc ggc aat gtt agc ggc		1392
	His Gln Pro Asn Thr Thr Phe Tyr Asp Tyr Thr Gly Asn Val Ser Gly		
	420	425	430
	430		
	acg gtg acc acc aat gcg cag ggc tat ggc agc ttc ccg gtc act atg		1440
35	Thr Val Thr Thr Asn Ala Gln Gly Tyr Gly Ser Phe Pro Val Thr Met		
	435	440	445
	445		
	acg gaa agt acc ggt tgg tca gtc tgg gta cca caa tcc aat ggt ggc		1488
	Thr Glu Ser Thr Gly Trp Ser Val Trp Val Pro Gln Ser Asn Gly Gly		

450 455 460 465

act cag ccg gga tcc att acc ctg cg<sup>g</sup> atg acc aag gat gtt ggc tat 1536  
Thr Gln Pro Gly Ser Ile Thr Leu Arg Met Thr Lys Asp Val Gly Tyr

5 470 475 480

ggc ttt tcg ttg ttc acc ggc agc agt g<sup>c</sup> gaa ctg acc aac tgg 1584  
Gly Phe Ser Leu Phe Phe Thr Gly Ser Ser Ala Glu Leu Thr Asn Trp

485 490 495

10 ggc ggc ggt att gaa ggc acc tgg aca tcc ggt aat gtc tgg gaa gtg 1632  
Gly Gly Gly Ile Glu Gly Thr Trp Thr Ser Gly Asn Val Trp Glu Val

500 505 510

15 acc atc ccg gat cc 1646  
Thr Ile Pro Asp

515

20 <210> 2  
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<212> PRT  
<213> *Bacillus* sp.

25 <220>  
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<222> (1)...(31)

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-30 -25 -20

Ala Ile Ala Leu Ser Ala Leu Ser Leu Ser Phe Gly Leu Gln Ala Ser  
-15 -10 -5 1

Glu Leu Pro Gln Ile Pro Pro Gln Gln Val Asn Asn Thr Met Tyr Gln

35 5 10 15

Ala Phe Tyr Trp Asp Ala Tyr Pro Gly Leu Trp Ala Asn Leu Pro Ala  
20 25 30

Met Ala Ala Pro Leu Ala Glu Arg Gly Ile Thr Ser Met Trp Leu Pro  
35 40 45

Pro Ala Ala Lys Gly Met Asn Gly Thr Phe Ser Val Gly Tyr Asp Val  
50 55 60 65

Tyr Asp Phe Trp Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Thr Ala  
70 75 80

5 Thr Arg Tyr Gly Thr Arg Gln Gln Leu Gln Gln Ala Leu Ser Ala Leu  
85 90 95

Asp Gln Leu Gly Ile Gln Ala Tyr Phe Asp Val Val Phe Asn His Arg  
100 105 110

Met Gly Ala Asp Ala Gln Glu Asn Ile Pro Gly Phe Gly Leu Ala Trp  
10 115 120 125

Thr Glu Tyr His Leu Gln Gly Arg Gln Ala His Tyr Thr Gln Gln Asn  
130 135 140 145

Trp Gly Tyr Leu Trp His Asp Phe Asp Trp Asn Trp Thr Ala Phe Asn  
150 155 160

15 Gly Ser Asp Asn Gln Leu Tyr Pro Gly Lys Trp Trp Gly Asn Thr Phe  
165 170 175

His Phe Pro Tyr Leu Met Gly Glu Asp Val Asp Tyr Asn Arg Phe Glu  
180 185 190

Val Gln Gln Glu Met Lys Ala Trp Gly Glu Trp Ile Ile Asn Ser Val  
20 195 200 205

Gly Phe Ser Gly Phe Arg Met Asp Ala Ile Ala His Val Asp Thr Asp  
210 215 220 225

Phe Thr Arg Asp Trp Ile Asn His Val Gln Trp Ala Thr Ser Glu Asp  
230 235 240

25 Val Phe Phe Val Ala Glu Ala Trp Val Ser Asp Ile Asn Gly Tyr Leu  
245 250 255

Asp Ala Val Asn Thr Pro His Leu Arg Ala Phe Asp Phe Asn Leu Arg  
260 265 270

Glu Asp Phe Val Ala Leu Ser Ser Gly Ser Lys Asp Met Arg Trp Trp  
30 275 280 285

Gly Gly Leu Val Asn Ser Gln His Arg Asp Arg Ala Val Thr Phe Val  
290 295 300 305

Asp Asn His Asp Thr Ser Arg Ala Gly Asn Pro Tyr Gly Met Pro Gln  
310 315 320

35 Val Ile Asn Tyr Lys Asn Gln Ala Tyr Ala Tyr Ile Leu Leu Arg Glu  
325 330 335

His Gly Val Pro Thr Val Phe Ala Arg Asp Tyr Asp Glu Phe Gly Met  
340 345 350

Ala Pro Thr Leu Asp Lys Leu Ile Glu Ala Arg Arg Tyr Phe Ala Tyr

355 360 365

Gly Pro Gly His Glu Tyr Ser Gly Asn Thr Glu Ala Val Tyr Ala Tyr  
370 375 380 385

Val Arg Glu Gly Leu Ser Thr Val Pro Gly Thr Gly Leu Val Met Leu  
5 390 395 400

Ile Ser Gly Arg Asn Trp Gly Gly Gln Gln Ser Phe Thr Ile Asn Ser  
405 410 415

His Gln Pro Asn Thr Thr Phe Tyr Asp Tyr Thr Gly Asn Val Ser Gly  
420 425 430

10 Thr Val Thr Asn Ala Gln Gly Tyr Ser Phe Pro Val Thr Met  
435 440 445

Thr Glu Ser Thr Gly Trp Ser Val Trp Val Pro Gln Ser Asn Gly Gly  
450 455 460 465

Thr Gln Pro Gly Ser Ile Thr Leu Arg Met Thr Lys Asp Val Gly Tyr  
15 470 475 480

Gly Phe Ser Leu Phe Phe Thr Gly Ser Ser Ala Glu Leu Thr Asn Trp  
485 490 495

Gly Gly Gly Ile Glu Gly Thr Trp Thr Ser Gly Asn Val Trp Glu Val  
500 505 510

20 Thr Ile Pro Asp  
515

<210> 3  
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25 <212> DNA  
<213> *Bacillus* sp.

<220>  
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30 <222> (1)...(1764)

<221> *sig\_peptide*  
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35 <221> *mat\_peptide*  
<222> (94)...(1764)

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	-30	-25	-20	
	gcc att gcc tta tcc gct ctc agt tta tcc ttt ggc ctg cag gcc agc			95
5	Ala Ile Ala Leu Ser Ala Leu Ser Leu Ser Phe Gly Leu Gln Ala Ser			
	-15	-10	-5	1
	gag tta cca caa att cca cca cag cag gtg aac aac acc atg tac cag			144
	Glu Leu Pro Gln Ile Pro Pro Gln Gln Val Asn Asn Thr Met Tyr Gln			
10	5	10	15	
	gca ttt tat tgg gat gcc tac cct ggc ctt tgg gcc aat tta ccg gcc			192
	Ala Phe Tyr Trp Asp Ala Tyr Pro Gly Leu Trp Ala Asn Leu Pro Ala			
	20	25	30	
15	atg gcg gcc cct ttg gcc gag cgt ggc att acc tcg atg tgg ttg ccg			240
	Met Ala Ala Pro Leu Ala Glu Arg Gly Ile Thr Ser Met Trp Leu Pro			
	35	40	45	
20	ccc gcc gcc aaa ggc atg aat ggt act ttc agt gtc ggt tac gat gta			288
	Pro Ala Ala Lys Gly Met Asn Gly Thr Phe Ser Val Gly Tyr Asp Val			
	50	55	60	65
	tac gat ttc tgg gat ctg ggc gag ttt aac caa aaa ggc acc acc gcc			336
25	Tyr Asp Phe Trp Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Thr Ala			
	70	75	80	
	acc cgt tac ggt act cgt cag cag ctg caa caa gca ctg agt gct ctg			384
	Thr Arg Tyr Gly Thr Arg Gln Gln Leu Gln Gln Ala Leu Ser Ala Leu			
30	85	90	95	
	gac caa ctg ggt att cag gcc tat ttt gat gtg gtg ttt aac cac cgc			432
	Asp Gln Leu Gly Ile Gln Ala Tyr Phe Asp Val Val Phe Asn His Arg			
	100	105	110	
35	atg ggc gcc gat gca cag gag aat att cct ggc ttt ggc ctg gcc tgg			480
	Met Gly Ala Asp Ala Gln Glu Asn Ile Pro Gly Phe Gly Leu Ala Trp			
	115	120	125	

acc gag tat cat ctg caa ggt cgt cag gcg cat tat acc cag caa aac 528  
Thr Glu Tyr His Leu Gln Gly Arg Gln Ala His Tyr Thr Gln Gln Asn  
130 135 140 145

5 tgg ggc tac ttg tgg cac gac ttt gac tgg aac tgg acc gcg ttt aat 576  
Trp Gly Tyr Leu Trp His Asp Phe Asp Trp Asn Trp Thr Ala Phe Asn  
150 155 160

10 ggc tcc gac aat cag ctc tac ccc ggc aaa tgg tgg ggc aat acc ttc 624  
Gly Ser Asp Asn Gln Leu Tyr Pro Gly Lys Trp Trp Gly Asn Thr Phe  
165 170 175

15 cac ttc cct tat ttg atg ggt gag gat gtc gat tac aac cgc ttt gaa 672  
His Phe Pro Tyr Leu Met Gly Glu Asp Val Asp Tyr Asn Arg Phe Glu  
180 185 190

20 gtg cag cag gaa atg aaa gcc tgg ggc gag tgg atc atc aac agc gtt 720  
Val Gln Gln Glu Met Lys Ala Trp Gly Glu Trp Ile Ile Asn Ser Val  
195 200 205

25 ggc ttt agc ggc ttt cgg atg gat gcc atc gcc cat gtc gat acc gat 768  
Gly Phe Ser Gly Phe Arg Met Asp Ala Ile Ala His Val Asp Thr Asp  
210 215 220 225

30 ttt acc cgt gac tgg atc aat cac gtg cag tgg gcc acc agt gag gat 816  
Phe Thr Arg Asp Trp Ile Asn His Val Gln Trp Ala Thr Ser Glu Asp  
230 235 240

35 gtg ttc ttt gtc gct gaa gcc tgg gtc agt gat atc aac ggc tat ctg 854  
Val Phe Phe Val Ala Glu Ala Trp Val Ser Asp Ile Asn Gly Tyr Leu  
245 250 255

40 gat gca gtc aat acg ccg cat ttg cgc gct ttt gat ttc aat ttg cgc 912  
Asp Ala Val Asn Thr Pro His Leu Arg Ala Phe Asp Phe Asn Leu Arg  
260 265 270

45 gaa gac ttc gtt gct tta agc agc ggt agc aaa gac atg cgt tgg tgg 950  
Glu Asp Phe Val Ala Leu Ser Ser Gly Ser Lys Asp Met Arg Trp Trp  
275 280 285

ggc ggt ctg gtc aat agc cag cac cgt gat cgg gcg gtc act ttt gtc 1008  
Gly Gly Leu Val Asn Ser Gln His Arg Asp Arg Ala Val Thr Phe Val  
290 295 300 305

5  
gat aac cac gat acc agc cgg gcc ggc aac cct tat ggc atg ccg cag 1056  
Asp Asn His Asp Thr Ser Arg Ala Gly Asn Pro Tyr Gly Met Pro Gln  
310 315 320

10  
gtg atc aac tac aag aac cag gcc tac gct tac att ctg ttg cgt gag 1104  
Val Ile Asn Tyr Lys Asn Gln Ala Tyr Ala Tyr Ile Leu Leu Arg Glu  
325 330 335

15  
cat ggg gtg ccg act gtg ttt gcc cgc gat tac gac gaa ttt ggc atg 1152  
His Gly Val Pro Thr Val Phe Ala Arg Asp Tyr Asp Glu Phe Gly Met  
340 345 350

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gcg cca acg ctg gat aaa ttg att gag gcg cgc cgc tac ttt gct tat 1200  
Ala Pro Thr Leu Asp Lys Leu Ile Glu Ala Arg Arg Tyr Phe Ala Tyr  
355 360 365

25  
ggt cct ggc cat gag tac tcc ggc aat acc gag gcc gtc tac gcc tat 1248  
Gly Pro Gly His Glu Tyr Ser Gly Asn Thr Glu Ala Val Tyr Ala Tyr  
370 375 380 385

gtg cgc gaa ggg ctt agc act gtg ccg ggt acc ggt ctg gtg atg ctg 1296  
Val Arg Glu Gly Leu Ser Thr Val Pro Gly Thr Gly Leu Val Met Leu  
390 395 400

30  
ata tcg ggt cga aac tgg ggt ggt cag cag tcg ttc acc atc aac agc 1344  
Ile Ser Gly Arg Asn Trp Gly Gly Gln Gln Ser Phe Thr Ile Asn Ser  
405 410 415

35  
cac cag ccg aat acc acc ttt tac gat tat acc ggc aat gtt agc ggc 1392  
His Gln Pro Asn Thr Thr Phe Tyr Asp Tyr Thr Gly Asn Val Ser Gly  
420 425 430

acg gtg acc acc aat gcg cag ggc tat ggc agc ttc ccg gtc act atg 1440  
Thr Val Thr Thr Asn Ala Gln Gly Tyr Gly Ser Phe Pro Val Thr Met

435 440 445

5 acg gaa agt acc ggt tgg tca gtc tgg gta cca caa tcc aat ggt ggc 1488  
Thr Glu Ser Thr Gly Trp Ser Val Trp Val Pro Gln Ser Asn Gly Gly  
450 455 460 465

10 act cag ccg gga tcc att acc ctg cgg atg acc aag gat gtt ggc tat 1536  
Thr Gln Pro Gly Ser Ile Thr Leu Arg Met Thr Lys Asp Val Gly Tyr  
470 475 480

15 ggc ttt tcg ttg ttc acc ggc agc agt gcg gaa ctg acc aac tgg 1584  
Gly Phe Ser Leu Phe Phe Thr Gly Ser Ser Ala Glu Leu Thr Asn Trp  
485 490 495

20 ggc ggc ggt att gaa ggc acc tgg aca tcc ggt aat gtc tgg gaa gtg 1632  
Gly Gly Gly Ile Glu Gly Thr Trp Thr Ser Gly Asn Val Trp Glu Val  
500 505 510

25 acc atc ccg gat ccg ggc aac ttt gaa tgg aaa acc cgt aaa ggc cca 1680  
Thr Ile Pro Asp Pro Gly Asn Phe Glu Trp Lys Thr Arg Lys Gly Pro  
515 520 525

30 acc ggt ggc agt ggt cag gac tgg gaa agt ggc agc aac cac aat cag 1728  
Thr Gly Gly Ser Gly Gln Asp Trp Glu Ser Gly Ser Asn His Asn Gln  
530 535 540 545

35 acc aat ttg cac ccc agt ttt aat ggt ggg ttt taa 1764  
Thr Asn Leu His Pro Ser Phe Asn Gly Gly Phe \*  
550 555

30 <210> 4  
<211> 587  
<212> PRT  
<213> *Bacillus* sp.

<220>  
<221> SIGNAL  
<222> (1)...(31)

&lt;400&gt; 4

Met Gln Asn Thr Ala Lys Asn Ser Ile Trp Gln Arg Val Arg His Ser  
-30 -25 -20

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-15 -10 -5 1

Glu Leu Pro Gln Ile Pro Pro Gln Gln Val Asn Asn Thr Met Tyr Gln  
5 10 15

Ala Phe Tyr Trp Asp Ala Tyr Pro Gly Leu Trp Ala Asn Leu Pro Ala  
10 20 25 30

Met Ala Ala Pro Leu Ala Glu Arg Gly Ile Thr Ser Met Trp Leu Pro  
35 40 45

Pro Ala Ala Lys Gly Met Asn Gly Thr Phe Ser Val Gly Tyr Asp Val  
50 55 60 65

15 Tyr Asp Phe Trp Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Thr Ala  
70 75 80

Thr Arg Tyr Gly Thr Arg Gln Gln Leu Gln Gln Ala Leu Ser Ala Leu  
85 90 95

Asp Gln Leu Gly Ile Gln Ala Tyr Phe Asp Val Val Phe Asn His Arg  
20 100 105 110

Met Gly Ala Asp Ala Gln Glu Asn Ile Pro Gly Phe Gly Leu Ala Trp  
115 120 125

Thr Glu Tyr His Leu Gln Gly Arg Gln Ala His Tyr Thr Gln Gln Asn  
130 135 140 145

25 Trp Gly Tyr Leu Trp His Asp Phe Asp Trp Asn Trp Thr Ala Phe Asn  
150 155 160

Gly Ser Asp Asn Gln Leu Tyr Pro Gly Lys Trp Trp Gly Asn Thr Phe  
165 170 175

His Phe Pro Tyr Leu Met Gly Glu Asp Val Asp Tyr Asn Arg Phe Glu  
30 180 185 190

Val Gln Gln Glu Met Lys Ala Trp Gly Glu Trp Ile Ile Asn Ser Val  
195 200 205

Gly Phe Ser Gly Phe Arg Met Asp Ala Ile Ala His Val Asp Thr Asp  
210 215 220 225

35 Phe Thr Arg Asp Trp Ile Asn His Val Gln Trp Ala Thr Ser Glu Asp  
230 235 240

Val Phe Phe Val Ala Glu Ala Trp Val Ser Asp Ile Asn Gly Tyr Leu  
245 250 255

Asp Ala Val Asn Thr Pro His Leu Arg Ala Phe Asp Phe Asn Leu Arg

260 265 270

Glu Asp Phe Val Ala Leu Ser Ser Gly Ser Lys Asp Met Arg Trp Trp  
275 280 285

Gly Gly Leu Val Asn Ser Gln His Arg Asp Arg Ala Val Thr Phe Val  
5 290 295 300 305

Asp Asn His Asp Thr Ser Arg Ala Gly Asn Pro Tyr Gly Met Pro Gln  
310 315 320

Val Ile Asn Tyr Lys Asn Gln Ala Tyr Ala Tyr Ile Leu Leu Arg Glu  
325 330 335

10 His Gly Val Pro Thr Val Phe Ala Arg Asp Tyr Asp Glu Phe Gly Met  
340 345 350

Ala Pro Thr Leu Asp Lys Leu Ile Glu Ala Arg Arg Tyr Phe Ala Tyr  
355 360 365

Gly Pro Gly His Glu Tyr Ser Gly Asn Thr Glu Ala Val Tyr Ala Tyr  
15 370 375 380 385

Val Arg Glu Gly Leu Ser Thr Val Pro Gly Thr Gly Leu Val Met Leu  
390 395 400

Ile Ser Gly Arg Asn Trp Gly Gly Gln Gln Ser Phe Thr Ile Asn Ser  
405 410 415

20 His Gln Pro Asn Thr Thr Phe Tyr Asp Tyr Thr Gly Asn Val Ser Gly  
420 425 430

Thr Val Thr Thr Asn Ala Gln Gly Tyr Ser Phe Pro Val Thr Met  
435 440 445

Thr Glu Ser Thr Gly Trp Ser Val Trp Val Pro Gln Ser Asn Gly Gly  
25 450 455 460 465

Thr Gln Pro Gly Ser Ile Thr Leu Arg Met Thr Lys Asp Val Gly Tyr  
470 475 480

Gly Phe Ser Leu Phe Phe Thr Gly Ser Ser Ala Glu Leu Thr Asn Trp  
485 490 495

30 Gly Gly Gly Ile Glu Gly Thr Trp Thr Ser Gly Asn Val Trp Glu Val  
500 505 510

Thr Ile Pro Asp Pro Gly Asn Phe Glu Trp Lys Thr Arg Lys Gly Pro  
515 520 525

Thr Gly Gly Ser Gly Gln Asp Trp Glu Ser Gly Ser Asn His Asn Gln  
35 530 535 540 545

Thr Asn Leu His Pro Ser Phe Asn Gly Gly Phe  
550 555